## SEQUENCE LISTING

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<110> LUCHE, Ralf M.
      WEI, Bo
<120> DSP-14 DUAL-SPECIFICITY PHOSPHATASE
<130> 200125.422US
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<150> 60/201,322
<151> 2000-05-02
<160> 16
<170> PatentIn Ver. 2.1
<210> 1
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<213> Homo sapiens
<400> 1
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cgcggccagg ccccggcaca cccagctgca gaaaggagag aaaatccctt ggctctaaaa 120
tgacatctgg agaagtgaag acaagcctca agaatgccta ctcatctgcc aagaggctgt 180
cgccgaagat ggaggaggaag aggactactg cacccctgga gcctttgagc 240
tggagcggct cttctggaag ggcagtcccc agtacaccca cgtcaacgag gtctggccca 300
agetetacat tggcgatgag gcgacggcgc tggaccgcta taggetgcag aaggeggggt 360
tcacgcacgt gctgaacgcg gcccacggcc gctggaacgt ggacactggg cccgactact 420
accgcgacat ggacatccag taccacggcg tggaggccga cgacctgccc accttcgacc 480
tcagtgtctt cttctacccg gcggcagcct tcatcgacag agcgctaagc gacgaccaca 540
gtaagateet ggtteaetge gteatgggee geageeggte agceaecetg gteetggeet 600
acctgatgat ccacaaggac atgaccctgg tggacgccat ccagcaagtg gccaagaacc 660
gctgcgtcct cccgaaccgg ggctttttga agcagctccg ggagctggac aagcagctgg 720
tgcagcagag gcgacggtcc cagcgccagg acggtgagga ggaggatggc agggagctgt 780
aggecegaet cacagggeca geagaggeae ttggggaeag aggggagagg cagaacatag 840
ccctggccta ggactccaga gaagggatgg tgaaaccgaa gctcgactct tccaaaccat 900
cttgttcaac ttccccatgt gtgctgggga cagggaggac ccagagctgc ccccgggcag 960
agctgagcgc tcagcctctc agcaaaatgg gagggacggg ctccccggct ctgggtcaca 1020
gaggagcatg ccacgctgca ccaagtctcc tgctttggtt ttgttttttt ggtgagaagg 1080
aagagggaaa aagattttta aaatgtgtag gcagtatgtt gtgattaaac gtttggcttt 1140
gtccaaaaaa aaaaaaaaaa aaaaa
                                                                  1165
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<211> 220
<212> PRT
<213> Homo sapiens
<400> 2
Met Thr Ser Gly Glu Val Lys Thr Ser Leu Lys Asn Ala Tyr Ser Ser
                  5
                                      10
                                                          15
Ala Lys Arg Leu Ser Pro Lys Met Glu Glu Glu Glu Glu Glu Glu Asp
             20
                                  25
Tyr Cys Thr Pro Gly Ala Phe Glu Leu Glu Arg Leu Phe Trp Lys Gly
         35
                              40
Ser Pro Gln Tyr Thr His Val Asn Glu Val Trp Pro Lys Leu Tyr Ile
     50
                          55
                                              60
Gly Asp Glu Ala Thr Ala Leu Asp Arg Tyr Arg Leu Gln Lys Ala Gly
 65
                     70
                                          75
Phe Thr His Val Leu Asn Ala Ala His Gly Arg Trp Asn Val Asp Thr
                 85
                                      90
Gly Pro Asp Tyr Tyr Arg Asp Met Asp Ile Gln Tyr His Gly Val Glu
            100
                                 105
                                                     110
Ala Asp Asp Leu Pro Thr Phe Asp Leu Ser Val Phe Phe Tyr Pro Ala
        115
                            120
                                                 125
Ala Ala Phe Ile Asp Arg Ala Leu Ser Asp Asp His Ser Lys Ile Leu
    130
                        135
                                             140
Val His Cys Val Met Gly Arg Ser Arg Ser Ala Thr Leu Val Leu Ala
145
                    150
                                         155
Tyr Leu Met Ile His Lys Asp Met Thr Leu Val Asp Ala Ile Gln Gln
                165
                                     170
Val Ala Lys Asn Arg Cys Val Leu Pro Asn Arg Gly Phe Leu Lys Gln
            180
                                185
                                                     190
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Arg Gln Asp Gly Glu Glu Glu Asp Gly Arg Glu Leu 210 215 220

Leu Arg Glu Leu Asp Lys Gln Leu Val Gln Gln Arg Arg Ser Gln

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<210> 3
<211> 19
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Peptide
<400> 3
Val His Cys Val Met Gly Arg Ser Arg Ser Ala Thr Leu Val Leu Ala
                                      10
                                                           15
Tyr Leu Met
<210> 4
<211> 24
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Peptide
Asn Gly Arg Val Leu Val His Cys Gln Ala Gly Ile Ser Arg Ser Gly
                  5
                                      10
                                                          15
Thr Asn Ile Leu Ala Tyr Leu Met
             20
<210> 5
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Nucleotide
      primer
<400> 5
tggcgtccac cagggtcatg tccttgtg
                                                                    28
<210> 6
<211> 28
<212> DNA
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<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Nucleotide
      primer
<400> 6
cacaaggaca tgaccctggt ggacgcca
                                                                    28
<210> 7
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Nucleotide
      primer
<400> 7
gccccagccg gtcagccacc ct
                                                                    22
<210> 8
<211> 170
<212> PRT
<213> Homo sapiens
<400> 8
Ser Asp Leu Asp Arg Asp Pro Asn Ser Ala Thr Asp Ser Asp Gly Ser
                                      10
Pro Leu Ser Asn Ser Gln Pro Ser Phe Pro Val Glu Ile Leu Pro Phe
             20
                                  25
                                                      30
Leu Tyr Leu Gly Cys Ala Lys Asp Ser Thr Asn Leu Asp Val Leu Glu
         35
                             40
Glu Phe Gly Ile Lys Tyr Ile Leu Asn Val Thr Pro Asn Leu Pro Asn
                         55
                                              60
Leu Phe Glu Asn Ala Gly Glu Phe Lys Tyr Lys Gln Ile Pro Ile Ser
 65
                     70
                                          75
Asp His Trp Ser Gln Asn Leu Ser Gln Phe Phe Pro Glu Ala Ile Ser
                 85
                                      90
Phe Ile Asp Glu Ala Arg Gly Lys Asn Cys Gly Val Leu Val His Cys
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100 105 110

Leu Ala Gly Ile Ser Arg Ser Val Thr Val Thr Val Ala Tyr Leu Met
115 120 125

Gln Lys Leu Asn Leu Ser Met Asn Asp Ala Tyr Asp Ile Val Lys Met 130 135 140

Lys Lys Ser Asn Ile Ser Pro Asn Phe Asn Phe Met Gly Gln Leu Leu 145 150 155 160

Asp Phe Glu Arg Thr Leu Gly Leu Ser Ser 165 170

<210> 9

<211> 168

<212> PRT

<213> Homo sapiens

<400> 9

Asp Arg Glu Leu Pro Ser Ser Ala Thr Glu Ser Asp Gly Ser Pro Val

1 5 10 15

Pro Ser Ser Gln Pro Ala Phe Pro Val Gln Ile Leu Pro Tyr Leu Tyr 20 25 30

Leu Gly Cys Ala Lys Asp Ser Thr Asn Leu Asp Val Leu Gly Lys Tyr
35 40 45

Gly Ile Lys Tyr Ile Leu Asn Val Thr Pro Asn Leu Pro Asn Ala Phe
50 55 60

Glu His Gly Gly Glu Phe Thr Tyr Lys Gln Ile Pro Ile Ser Asp His 65 70 75 80

Trp Ser Gln Asn Leu Ser Gln Phe Phe Pro Glu Ala Ile Ser Phe Ile 85 90 95

Asp Glu Ala Arg Ser Lys Lys Cys Gly Val Leu Val His Cys Leu Ala 100 105 110

Gly Ile Ser Arg Ser Val Thr Val Thr Val Ala Tyr Leu Met Gln Lys 115 120 125

Met Asn Leu Ser Leu Asn Asp Ala Tyr Asp Phe Val Lys Arg Lys Lys 130 135 140

Ser Asn Ile Ser Pro Asn Phe Asn Phe Met Gly Gln Leu Leu Asp Phe 145 150 155 160

Glu Arg Thr Leu Gly Leu Ser Ser 165

<210> 10

<211> 157

<212> PRT

<213> Homo sapiens

<400> 10

Gly Ala Thr Pro Pro Pro Val Gly Leu Arg Ala Ser Phe Pro Val Gln
1 5 10 15

Ile Leu Pro Asn Leu Tyr Leu Gly Ser Ala Arg Asp Ser Ala Asn Leu
20 25 30

Glu Ser Leu Ala Lys Leu Gly Ile Arg Tyr Ile Leu Asn Val Thr Pro  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Asn Leu Pro Asn Phe Phe Glu Lys Asn Gly Asp Phe His Tyr Lys Gln 50 55 60

Ile Pro Ile Ser Asp His Trp Ser Gln Asn Leu Ser Arg Phe Phe Pro 65 70 75 80

Glu Ala Ile Glu Phe Ile Asp Glu Ala Leu Ser Gln Asn Cys Gly Val\$85\$ 90 95

Leu Val His Cys Leu Ala Gly Val Ser Arg Ser Val Thr Val Thr Val 100 105 110

Ala Tyr Leu Met Gln Lys Leu His Leu Ser Leu Asn Asp Ala Tyr Asp 115 120 125

Leu Val Lys Arg Lys Lys Ser Asn Ile Ser Pro Asn Phe Asn Phe Met 130 135 140

<210> 11

<211> 170

<212> PRT

<213> Homo sapiens

<400> 11															
Gly 1	Leu	Cys	Glu	Gly 5	Lys	Pro	Ala	Ala	Leu 10	Leu	Pro	Met	Ser	Leu 15	Ser
Gln	Pro	Cys	Leu 20	Pro	Val	Pro	Ser	Val 25	Gly	Leu	Thr	Arg	Ile 30	Leu	Pro
His	Leu	Tyr 35	Leu	Gly	Ser	Gln	Lys 40	Asp	Val	Leu	Asn	Lys 45	Asp	Leu	Met
Thr	Gln 50	Asn	Gly	Ile	Ser	Tyr 55	Val	Leu	Asn	Ala	Ser 60	Asn	Ser	Cys	Pro
Lys 65	Pro	Asp	Phe	Ile	Cys 70	Glu	Ser	Arg	Phe	Met 75	Arg	Val	Pro	Ile	Asn 80
Asp	Asn	Tyr	Cys	Glu 85	Lys	Leu	Leu	Pro	Trp 90	Leu	Asp	Lys	Ser	Ile 95	Glu
Phe	I <b>l</b> e	Asp	Lys 100	Ala	Lys	Leu	Ser	Ser 105	Cys	Gln	Val	Ile	Val 110	His	Cys
Leu	Ala	Gly 115	Ile	Ser	Arg	Ser	Ala 120	Thr	Ile	Ala	Ile	Ala 125	Tyr	Ile	Met
Lys	Thr 130	Met	Gly	Met	Ser	Ser 135	Asp	Asp	Ala	Tyr	Arg 140	Phe	Val	Lys	Asp
Arg 145	Arg	Pro	Ser	Ile	Ser 150	Pro	Asn	Phe	Asn	Phe 155	Leu	Gly	Gln	Leu	Leu 160
Glu	Tyr	Glu	Arg	Thr 165	Leu	Lys	Leu	Leu	Ala 170						
<210	> 12														

<210> 12 <211> 168 <212> PRT <213> Homo sapiens

<400> 12

Pro Ala Gln Ala Leu Pro Pro Ala Gly Ala Glu Asn Ser Asn Ser Asp 1 5 10 15

Pro Arg Val Pro Ile Tyr Asp Gln Gly Gly Pro Val Glu Ile Leu Pro 20 25 30

Tyr Leu Tyr Leu Gly Ser Cys Asn His Ser Ser Asp Leu Gln Gly Leu 35 40 45

Gln Ala Cys Gly Ile Thr Ala Val Leu Asn Val Ser Ala Ser Cys Pro 50 55 60

Asn His Phe Glu Gly Leu Phe His Tyr Lys Ser Ile Pro Val Glu Asp
65 70 75 80

Asn Gln Met Val Glu Ile Ser Ala Trp Phe Gln Glu Ala Ile Ser Phe
85 90 95

Ile Asp Ser Val Lys Asn Ser Gly Gly Arg Val Leu Val His Cys Gln
100 105 110

Ala Gly Ile Ser Arg Ser Ala Thr Ile Cys Leu Ala Tyr Leu Ile Gln
115 120 125

Ser His Arg Val Arg Leu Asp Glu Ala Phe Asp Phe Val Lys Gln Arg 130 135 140

Arg Gly Val Ile Ser Pro Asn Phe Ser Phe Met Gly Gln Leu Leu Gln 145 150 155 160

Leu Glu Thr Gln Val Leu Cys His 165

<210> 13

<211> 169

<212> PRT

<213> Homo sapiens

<400> 13

Pro Leu Ser Thr Ser Val Pro Asp Ser Ala Glu Ser Gly Cys Ser Ser 1 5 10 15

Cys Ser Thr Pro Leu Tyr Asp Gln Gly Gly Pro Val Glu Ile Leu Pro
20 25 30

Phe Leu Tyr Leu Gly Ser Ala Tyr His Ala Ser Arg Lys Asp Met Leu 35 40 45

Asp Ala Leu Gly Ile Thr Ala Leu Ile Asn Val Ser Ala Asn Cys Pro 50 55 60

Asn His Phe Glu Gly His Tyr Gln Tyr Lys Ser Ile Pro Val Glu Asp 65 70 75 80

Asn His Lys Ala Asp Ile Ser Ser Trp Phe Asn Glu Ala Ile Asp Phe
85 90 95

Ile Asp Ser Ile Lys Asn Ala Gly Gly Arg Val Phe Val His Cys Gln
100 105 110

Ala Gly Ile Ser Arg Ser Ala Thr Ile Cys Leu Ala Tyr Leu Met Arg 115 120 125

Thr Asn Arg Val Lys Leu Asp Glu Ala Phe Glu Phe Val Lys Gln Arg 130 135 140

Phe Glu Ser Gln Val Leu Ala Pro His 165

<210> 14

<211> 169

<212> PRT

<213> Homo sapiens

<400> 14

Pro Val Pro Pro Ser Ala Thr Glu Pro Leu Asp Leu Gly Cys Ser Ser 1 5 10 15

Cys Gly Thr Pro Leu His Asp Gln Gly Gly Pro Val Glu Ile Leu Pro 20 25 30

Phe Leu Tyr Leu Gly Ser Ala Tyr His Ala Ala Arg Arg Asp Met Leu 35 40 45

Asp Ala Leu Gly Ile Thr Ala Leu Leu Asn Val Ser Ser Asp Cys Pro 50 55 60

Asn His Phe Glu Gly His Tyr Gln Tyr Lys Cys Ile Pro Val Glu Asp
65 70 75 80

Asn His Lys Ala Asp Ile Ser Ser Trp Phe Met Glu Ala Ile Glu Tyr 85 90 95

Ile Asp Ala Val Lys Asp Cys Arg Gly Arg Val Leu Val His Cys Gln
100 105 110

Ala Gly Ile Ser Arg Ser Ala Thr Ile Cys Leu Ala Tyr Leu Met Met

115 120 125

Lys Lys Arg Val Arg Leu Glu Glu Ala Phe Glu Phe Val Lys Gln Arg 130 135 140

Phe Glu Ser Gln Val Leu Ala Thr Ser 165

<210> 15

<211> 171

<212> PRT

<213> Homo sapiens

<400> 15

Ser Glu Arg Ala Leu Ile Ser Gln Cys Gly Lys Pro Val Val Asn Val 1 5 10 15

Ser Tyr Arg Pro Ala Tyr Asp Gln Gly Gly Pro Val Glu Ile Leu Pro 20 25 30

Phe Leu Tyr Leu Gly Ser Ala Tyr His Ala Ser Lys Cys Glu Phe Leu 35 40 45

Ala Asn Leu His Ile Thr Ala Leu Leu Asn Val Ser Arg Arg Thr Ser 50 55 60

Glu Ala Cys Met Thr His Leu His Tyr Lys Trp Ile Pro Val Glu Asp
65 70 75 80

Ser His Thr Ala Asp Ile Ser Ser His Phe Gln Glu Ala Ile Asp Phe 85 90 95

Ile Asp Cys Val Arg Glu Lys Gly Gly Lys Val Leu Val His Cys Glu
100 105 110

Ala Gly Ile Ser Arg Ser Pro Thr Ile Cys Met Ala Tyr Leu Met Lys 115 120 125

Thr Lys Gln Phe Arg Leu Lys Glu Ala Phe Asp Tyr Ile Lys Gln Arg 130 135 140

Arg Ser Met Val Ser Pro Asn Phe Gly Phe Met Gly Gln Leu Leu Gln 145 150 155 160 Tyr Glu Ser Glu Ile Leu Pro Ser Thr Pro Asn 

<210> 16

<211> 180

<212> PRT

<213> Homo sapiens

<400> 16

Ser Gly Ser Phe Glu Leu Ser Val Gln Asp Leu Asn Asp Leu Leu Ser 

Asp Gly Ser Gly Cys Tyr Ser Leu Pro Ser Gln Pro Cys Asn Glu Val 

Thr Pro Arg Ile Tyr Val Gly Asn Ala Ser Val Ala Gln Asp Ile Pro 

Lys Leu Gln Lys Leu Gly Ile Thr His Val Leu Asn Ala Ala Glu Gly 

Arg Ser Phe Met His Val Asn Thr Asn Ala Asn Phe Tyr Lys Asp Ser 

Gly Ile Thr Tyr Leu Gly Ile Lys Ala Asn Asp Thr Gln Glu Phe Asn 

Leu Ser Ala Tyr Phe Glu Arg Ala Ala Asp Phe Ile Asp Gln Ala Leu 

Ala Gln Lys Asn Gly Arg Val Leu Val His Cys Arg Glu Gly Tyr Ser 

Arg Ser Pro Thr Leu Val Ile Ala Tyr Leu Met Met Arg Gln Lys Met 

Asp Val Lys Ser Ala Leu Ser Ile Val Arg Gln Asn Arg Glu Ile Gly 

Pro Asn Asp Gly Phe Leu Ala Gln Leu Cys Gln Leu Asn Asp Arg Leu 

Ala Lys Glu Gly